SEQUENCE LISTING

<110> Friddle, Carl Johan
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Scoville, John
Walke, D. Wade

<120> Novel Human Secreted Signal Proteins and Polynucleotides Encoding the Same

<130> LEX-0198-USA

<150> US 60/216,384
<151> 2000-07-07

<150> US 60/219,890
<151> 2000-07-21

<150> US 60/230,609
<151> 2000-09-06

<160> 24

<170> FastSEQ for Windows Version 4.0

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<212> DNA
<213> homo sapiens

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caacccgact tccgcgccat cggtgacttc ctcaaggaca agtacgacag cgcctcggag 720
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<211> 423
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<213> homo sapiens

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            20   25   30
Ser Ser Leu Gly Ser Gln Pro Ile Leu Cys Ala Ser Ile Pro Gly Leu
        35   40   45
Val Pro Lys Gln Leu Arg Phe Cys Arg Asn Tyr Val Glu Ile Met Pro
    50   55   60
Ser Val Ala Glu Gly Ile Lys Ile Gly Ile Gln Glu Cys Gln His Gln
65   70   75   80
Phe Arg Gly Arg Arg Trp Asn Cys Thr Thr Val His Asp Ser Leu Ala
                85   90   95
Ile Phe Gly Pro Val Leu Asp Lys Ala Thr Arg Glu Ser Ala Phe Val
            100   105   110
His Ala Ile Ala Ser Ala Gly Val Ala Phe Ala Val Thr Arg Ser Cys
        115   120   125
Ala Glu Gly Thr Ala Ala Ile Cys Gly Cys Ser Ser Arg His Gln Gly
    130   135   140
Ser Pro Gly Lys Gly Trp Lys Trp Gly Gly Cys Ser Glu Asp Ile Glu
145   150   155   160
Phe Gly Gly Met Val Ser Arg Glu Phe Ala Asp Ala Arg Glu Asn Arg
                165   170   175
Pro Asp Ala Arg Ser Ala Met Asn Arg His Asn Asn Glu Ala Gly Arg
            180   185   190
Gln Ala Ile Ala Ser His Met His Leu Lys Cys Lys Cys His Gly Leu
        195   200   205
Ser Gly Ser Cys Glu Val Lys Thr Cys Trp Trp Ser Gln Pro Asp Phe
    210   215   220
Arg Ala Ile Gly Asp Phe Leu Lys Asp Lys Tyr Asp Ser Ala Ser Glu
225   230   235   240
Met Val Val Glu Lys His Arg Glu Ser Arg Gly Trp Val Glu Thr Leu
                245   250   255
Arg Pro Arg Tyr Thr Tyr Phe Lys Val Pro Thr Glu Arg Asp Leu Val
            260   265   270
Tyr Tyr Glu Ala Ser Pro Asn Phe Cys Glu Pro Asn Pro Glu Thr Gly
        275   280   285
Ser Phe Gly Thr Arg Asp Arg Thr Cys Asn Val Ser Ser His Gly Ile
    290   295   300
Asp Gly Cys Asp Leu Leu Cys Cys Gly Arg Gly His Asn Ala Arg Ala
305   310   315   320
Glu Arg Arg Arg Glu Lys Cys Arg Cys Val Phe His Trp Cys Cys Tyr
                325   330   335
Val Ser Cys Gln Glu Cys Thr Arg Val Tyr Asp Val His Thr Cys Lys
            340   345   350
Asp Gly Cys Leu Arg Thr Gly His Ser Gly Pro Cys Arg Ser Leu Ala
        355   360   365
Trp Ile Trp Ser Pro Gly Ser Gln Gly His Asp Leu Leu Glu Gln Leu
    370   375   380
Pro Arg Ser Gly Gly Leu Gly Gln Cys Ser Ser Leu Gln Asn Trp Thr
385   390   395   400
Ala Val Ser Gly Cys Leu Arg Asp His Leu Gly Gly Leu Pro Gly Gly

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<210> 3
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<212> DNA
<213> homo sapiens

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ctgtgtgcca gcatcccggg cctggtcccc aagcagctcc gcttctgcag gaactacgtg     180
gagatcatgc ccagcgtggc cgagggcatc aagattggca tccaggagtg ccagcaccag     240
ttccgcggcc gccggtggaa ctgcaccacc gtccacgaca gcctggccat cttcgggccc     300
gtgctggaca aagctaccag ggagtcggcc tttgtccacg ccattgcctc agccggtgtg     360
gcctttgcag tgacacgctc atgtgcagaa ggcacggccg ccatctgtgg ctgcagcagc     420
cgccaccagg gctcaccagg caagggctgg aagtggggtg gctgtagcga ggacatcgag     480
tttggtggga tggtgtctcg ggagttcgcc gacgcccggg agaaccggcc agatgcccgc     540
tcagccatga accgccacaa caacgaggct gggcgccagg ccatcgccag ccacatgcac     600
ctcaagtgca agtgccacgg gctgtcgggc agctgcgagg tgaagacatg ctggtggtcg     660
caacccgact tccgcgccat cggtgacttc ctcaaggaca agtacgacag cgcctcggag     720
atggtggtgg agaagcaccg ggagtcccgc ggctgggtgg agaccctgcg gccgcgctac     780
acctacttca aggtgcccac ggagcgcgac ctggtctact acgaggcctc gcccaacttc     840
tgcgagccca accctgagac gggctccttc ggcacgcgcg accgcacctg caacgtcagc     900
tcgcacggca tcgacggctg cgacctgctg tgctgcggcc gcggccacaa cgcgcgagcg     960
gagcggcgcc gggagaagtg ccgctgcgtg ttccactggt gctgctacgt cagctgccag    1020
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<210> 4
<211> 352
<212> PRT
<213> homo sapiens

<400> 4

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            20                  25                  30
Ser Ser Leu Gly Ser Gln Pro Ile Leu Cys Ala Ser Ile Pro Gly Leu
        35                  40                  45
Val Pro Lys Gln Leu Arg Phe Cys Arg Asn Tyr Val Glu Ile Met Pro
    50                  55                  60
Ser Val Ala Glu Gly Ile Lys Ile Gly Ile Gln Glu Cys Gln His Gln
65                  70                  75                  80
Phe Arg Gly Arg Arg Trp Asn Cys Thr Thr Val His Asp Ser Leu Ala
                85                  90                  95
Ile Phe Gly Pro Val Leu Asp Lys Ala Thr Arg Glu Ser Ala Phe Val
            100                 105                 110
His Ala Ile Ala Ser Ala Gly Val Ala Phe Ala Val Thr Arg Ser Cys
        115                 120                 125
Ala Glu Gly Thr Ala Ala Ile Cys Gly Cys Ser Ser Arg His Gln Gly
    130                 135                 140
Ser Pro Gly Lys Gly Trp Lys Trp Gly Gly Cys Ser Glu Asp Ile Glu
145                 150                 155                 160
Phe Gly Gly Met Val Ser Arg Glu Phe Ala Asp Ala Arg Glu Asn Arg
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                165                 170                 175
Pro Asp Ala Arg Ser Ala Met Asn Arg His Asn Asn Glu Ala Gly Arg
            180                 185                 190
Gln Ala Ile Ala Ser His Met His Leu Lys Cys Lys Cys His Gly Leu
        195                 200                 205
Ser Gly Ser Cys Glu Val Lys Thr Cys Trp Trp Ser Gln Pro Asp Phe
    210                 215                 220
Arg Ala Ile Gly Asp Phe Leu Lys Asp Lys Tyr Asp Ser Ala Ser Glu
225                 230                 235                 240
Met Val Val Glu Lys His Arg Glu Ser Arg Gly Trp Val Glu Thr Leu
                245                 250                 255
Arg Pro Arg Tyr Thr Tyr Phe Lys Val Pro Thr Glu Arg Asp Leu Val
            260                 265                 270
Tyr Tyr Glu Ala Ser Pro Asn Phe Cys Glu Pro Asn Pro Glu Thr Gly
        275                 280                 285
Ser Phe Gly Thr Arg Asp Arg Thr Cys Asn Val Ser Ser His Gly Ile
    290                 295                 300
Asp Gly Cys Asp Leu Leu Cys Cys Gly Arg Gly His Asn Ala Arg Ala
305                 310                 315                 320
Glu Arg Arg Arg Glu Lys Cys Arg Cys Val Phe His Trp Cys Cys Tyr
                325                 330                 335
Val Ser Cys Gln Glu Cys Thr Arg Val Tyr Asp Val His Thr Cys Lys
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<210> 5
<211> 1505
<212> DNA
<213> homo sapiens

<400> 5

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ggccctccgc gccctctcgc gcggcgatgg ccccactcgg atacttctta ctcctctgca     180
gcctgaagca ggctctgggc agctacccga tctggtggtc gctggctgtt gggccacagt     240
attcctccct gggctcgcag cccatcctgt gtgccagcat cccgggcctg gtccccaagc     300
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acgacagcct ggccatcttc gggcccgtgc tggacaaagc taccagggag tcggcctttg     480
tccacgccat tgcctcagcc ggtgtggcct ttgcagtgac acgctcatgt gcagaaggca     540
cggccgccat ctgtggctgc agcagccgcc accagggctc accaggcaag ggctggaagt     600
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aggacaagta cgacagcgcc tcggagatgg tggtggagaa gcaccgggag tcccgcggct     900
gggtggagac cctgcggccg cgctacacct acttcaaggt gcccacggag cgcgacctgg     960
tctactacga ggcctcgccc aacttctgcg agcccaaccc tgagacgggc tccttcggca    1020
cgcgcgaccg cacctgcaac gtcagctcgc acggcatcga cggctgcgac ctgctgtgct    1080
gcggccgcgg ccacaacgcg cgagcggagc ggcgccggga gaagtgccgc tgcgtgttcc    1140
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gccctggatc ccaggggcat gacctcctgg agcaacttcc aaggtctggg ggactaggtc    1320
agtgttcctc cctgcagaac tggactgctg tcagtggctg ccttcgggat catctgggag    1380
ggctttcagg aggaggtgag catgggggaca cttctaggg ctccaacatc ctcctgtaat    1440
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ccctc                                                               1505
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<210> 6
<211> 1110
<212> DNA
<213> homo sapiens

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aggtcagtga acaatttcct gataacaggt cccaaggcct atctgaccta cacgactagt 180
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ggaggccatg gctggatctg gggaggctgc agcgacaatg tggaatttgg ggaaaggatc 480
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<210> 7
<211> 369
<212> PRT
<213> homo sapiens

<400> 7
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Cys Leu Thr Phe Ser Leu Phe Gly Arg Ser Val Asn Asn Phe Leu Ile
35 40 45
Thr Gly Pro Lys Ala Tyr Leu Thr Tyr Thr Thr Ser Val Ala Leu Gly
50 55 60
Ala Gln Ser Gly Ile Glu Glu Cys Lys Phe Gln Phe Ala Trp Glu Arg
65 70 75 80
Trp Asn Cys Pro Glu Asn Ala Leu Gln Leu Ser Thr His Asn Arg Leu
85 90 95
Arg Ser Ala Thr Arg Glu Thr Ser Phe Ile His Ala Ile Ser Ser Ala
100 105 110
Gly Val Met Tyr Ile Ile Thr Lys Asn Cys Ser Met Gly Asp Phe Glu
115 120 125
Asn Cys Gly Cys Asp Gly Ser Asn Asn Gly Lys Thr Gly Gly His Gly
130 135 140
Trp Ile Trp Gly Gly Cys Ser Asp Asn Val Glu Phe Gly Glu Arg Ile
145 150 155 160
Ser Lys Leu Phe Val Asp Ser Leu Glu Lys Gly Lys Asp Ala Arg Ala
165 170 175
Leu Met Asn Leu His Asn Asn Arg Ala Gly Arg Leu Ala Val Arg Ala
180 185 190

Thr Met Lys Arg Thr Cys Lys Cys His Gly Ile Ser Gly Ser Cys Ser
195 200 205
Ile Gln Thr Cys Trp Leu Gln Leu Ala Glu Phe Arg Glu Met Gly Asp
210 215 220
Tyr Leu Lys Ala Lys Tyr Asp Gln Ala Leu Lys Ile Glu Met Asp Lys
225 230 235 240
Arg Gln Leu Arg Ala Gly Asn Ser Ala Glu Gly His Trp Val Pro Ala
245 250 255
Glu Ala Phe Leu Pro Ser Ala Glu Ala Glu Leu Ile Phe Leu Glu Glu
260 265 270
Ser Pro Asp Tyr Cys Thr Cys Asn Ser Ser Leu Gly Ile Tyr Gly Thr
275 280 285
Glu Gly Arg Glu Cys Leu Gln Asn Ser His Asn Thr Ser Arg Trp Glu
290 295 300
Arg Arg Ser Cys Gly Arg Leu Cys Thr Glu Cys Gly Leu Gln Val Glu
305 310 315 320
Glu Arg Lys Thr Glu Val Ile Ser Ser Cys Asn Cys Lys Phe Gln Trp
325 330 335
Cys Cys Thr Val Lys Cys Asp Gln Cys Arg His Val Val Ser Lys Tyr
340 345 350
Tyr Cys Ala Arg Ser Pro Gly Ser Ala Gln Ser Leu Gly Lys Gly Ser
355 360 365
Ala

<210> 8
<211> 1056
<212> DNA
<213> homo sapiens

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actagtgtgg ccttgggtgc ccagagtggc atcgaggagt gcaagttcca gtttgcttgg 180
gaacgctgga actgccctga aaatgctctt cagctctcca cccacaacag gctgagaagt 240
gctaccagag agacttcctt catacatgct atcagctctg ctggagtcat gtacatcatc 300
accaagaact gtagcatggg tgacttcgaa aactgtggct gtgatgggtc aaacaatgga 360
aaaacaggag gccatggctg gatctgggga ggctgcagcg acaatgtgga atttggggaa 420
aggatctcca aactctttgt ggacagtttg gagaagggga aggatgccag agccctgatg 480
aatcttcaca acaacagggc cggcagactg gcagtgagag ccaccatgaa aaggacatgc 540
aaatgtcatg gcatctctgg gagctgcagc atacagacat gctggctgca gctggctgaa 600
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gataagcggc agctgagagc tgggaacagc gccgagggcc actgggtgcc cgctgaggcc 720
ttccttccta gcgcagaggc ggaactgatc tttttagagg aatcaccaga ttactgtacc 780
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<210> 9
<211> 351
<212> PRT
<213> homo sapiens

<400> 9

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            20                  25                  30
Pro Lys Ala Tyr Leu Thr Tyr Thr Thr Ser Val Ala Leu Gly Ala Gln
        35                  40                  45
Ser Gly Ile Glu Glu Cys Lys Phe Gln Phe Ala Trp Glu Arg Trp Asn
    50                  55                  60
Cys Pro Glu Asn Ala Leu Gln Leu Ser Thr His Asn Arg Leu Arg Ser
65                  70                  75                  80
Ala Thr Arg Glu Thr Ser Phe Ile His Ala Ile Ser Ser Ala Gly Val
                85                  90                  95
Met Tyr Ile Ile Thr Lys Asn Cys Ser Met Gly Asp Phe Glu Asn Cys
            100                 105                 110
Gly Cys Asp Gly Ser Asn Asn Gly Lys Thr Gly Gly His Gly Trp Ile
        115                 120                 125
Trp Gly Gly Cys Ser Asp Asn Val Glu Phe Gly Glu Arg Ile Ser Lys
    130                 135                 140
Leu Phe Val Asp Ser Leu Glu Lys Gly Lys Asp Ala Arg Ala Leu Met
145                 150                 155                 160
Asn Leu His Asn Asn Arg Ala Gly Arg Leu Ala Val Arg Ala Thr Met
                165                 170                 175
Lys Arg Thr Cys Lys Cys His Gly Ile Ser Gly Ser Cys Ser Ile Gln
            180                 185                 190
Thr Cys Trp Leu Gln Leu Ala Glu Phe Arg Glu Met Gly Asp Tyr Leu
        195                 200                 205
Lys Ala Lys Tyr Asp Gln Ala Leu Lys Ile Glu Met Asp Lys Arg Gln
    210                 215                 220
Leu Arg Ala Gly Asn Ser Ala Glu Gly His Trp Val Pro Ala Glu Ala
225                 230                 235                 240
Phe Leu Pro Ser Ala Glu Ala Glu Leu Ile Phe Leu Glu Glu Ser Pro
                245                 250                 255
Asp Tyr Cys Thr Cys Asn Ser Ser Leu Gly Ile Tyr Gly Thr Glu Gly
            260                 265                 270
Arg Glu Cys Leu Gln Asn Ser His Asn Thr Ser Arg Trp Glu Arg Arg
        275                 280                 285
Ser Cys Gly Arg Leu Cys Thr Glu Cys Gly Leu Gln Val Glu Glu Arg
    290                 295                 300
Lys Thr Glu Val Ile Ser Ser Cys Asn Cys Lys Phe Gln Trp Cys Cys
305                 310                 315                 320
Thr Val Lys Cys Asp Gln Cys Arg His Val Val Ser Lys Tyr Tyr Cys
                325                 330                 335
Ala Arg Ser Pro Gly Ser Ala Gln Ser Leu Gly Lys Gly Ser Ala
            340                 345                 350
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<210> 10
<211> 768
<212> DNA
<213> homo sapiens

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gaatttgggg aaaggatctc caaactcttt gtggacagtt tggagaaggg gaaggatgcc 180
agagccctga tgaatcttca caacaacagg gccggcagac tggcagtgag agccaccatg 240
aaaaggacat gcaaatgtca tggcatctct gggagctgca gcatacagac atgctggctg 300

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cagctggctg aattccggga gatgggagac tacctaaagg ccaagtatga ccaggcgctg     360
aaaattgaaa tggataagcg gcagctgaga gctgggaaca gcgccgaggg ccactgggtg     420
cccgctgagg ccttccttcc tagcgcagag gcggaactga tctttttaga ggaatcacca     480
gattactgta cctgcaattc cagcctgggc atctatggca cagagggtcg tgagtgccta     540
cagaacagcc acaacacatc caggtgggag cgacgtagct gtgggcgcct gtgcactgag     600
tgtgggctgc aggtggaaga gaggaaaact gaggtcataa gcagctgtaa ctgcaaattc     660
cagtggtgct gtacggtcaa gtgtgaccag tgtaggcatg tggtgagcaa gtattactgc     720
gcacgctccc caggcagtgc ccagtccctg ggtaagggca gtgcctga                 768

<210> 11
<211> 255
<212> PRT
<213> homo sapiens

<400> 11
Met Tyr Ile Ile Thr Lys Asn Cys Ser Met Gly Asp Phe Glu Asn Cys
 1               5                  10                  15
Gly Cys Asp Gly Ser Asn Asn Gly Lys Thr Gly Gly His Gly Trp Ile
            20                  25                  30
Trp Gly Gly Cys Ser Asp Asn Val Glu Phe Gly Glu Arg Ile Ser Lys
        35                  40                  45
Leu Phe Val Asp Ser Leu Glu Lys Gly Lys Asp Ala Arg Ala Leu Met
    50                  55                  60
Asn Leu His Asn Asn Arg Ala Gly Arg Leu Ala Val Arg Ala Thr Met
65                  70                  75                  80
Lys Arg Thr Cys Lys Cys His Gly Ile Ser Gly Ser Cys Ser Ile Gln
                85                  90                  95
Thr Cys Trp Leu Gln Leu Ala Glu Phe Arg Glu Met Gly Asp Tyr Leu
            100                 105                 110
Lys Ala Lys Tyr Asp Gln Ala Leu Lys Ile Glu Met Asp Lys Arg Gln
        115                 120                 125
Leu Arg Ala Gly Asn Ser Ala Glu Gly His Trp Val Pro Ala Glu Ala
    130                 135                 140
Phe Leu Pro Ser Ala Glu Ala Glu Leu Ile Phe Leu Glu Glu Ser Pro
145                 150                 155                 160
Asp Tyr Cys Thr Cys Asn Ser Ser Leu Gly Ile Tyr Gly Thr Glu Gly
                165                 170                 175
Arg Glu Cys Leu Gln Asn Ser His Asn Thr Ser Arg Trp Glu Arg Arg
            180                 185                 190
Ser Cys Gly Arg Leu Cys Thr Glu Cys Gly Leu Gln Val Glu Glu Arg
        195                 200                 205
Lys Thr Glu Val Ile Ser Ser Cys Asn Cys Lys Phe Gln Trp Cys Cys
    210                 215                 220
Thr Val Lys Cys Asp Gln Cys Arg His Val Val Ser Lys Tyr Tyr Cys
225                 230                 235                 240
Ala Arg Ser Pro Gly Ser Ala Gln Ser Leu Gly Lys Gly Ser Ala
                245                 250                 255

<210> 12
<211> 105
<212> DNA
<213> homo sapiens

<400> 12
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tctgcctggt cagtgaacaa tttcctgata acagggccca tctga                     105
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<210> 13
<211> 34
<212> PRT
<213> homo sapiens

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Ala Phe Ser Ala Ser Ala Trp Ser Val Asn Asn Phe Leu Ile Thr Gly
20 25 30
Pro Ile

<210> 14
<211> 72
<212> DNA
<213> homo sapiens

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<210> 15
<211> 23
<212> PRT
<213> homo sapiens

<400> 15
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<210> 16
<211> 111
<212> DNA
<213> homo sapiens

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<210> 17
<211> 36
<212> PRT
<213> homo sapiens

<400> 17
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Pro Lys Val Gly
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<210> 18

<211> 1056
<212> DNA
<213> homo sapiens

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<211> 351
<212> PRT
<213> homo sapiens

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Pro Lys Ala Tyr Leu Thr Tyr Thr Thr Ser Val Ala Leu Gly Ala Gln
        35   40   45
Ser Gly Ile Glu Glu Cys Lys Phe Gln Phe Ala Trp Glu Arg Trp Asn
    50   55   60
Cys Pro Glu Asn Ala Leu Gln Leu Ser Thr His Asn Arg Leu Arg Ser
65   70   75   80
Ala Thr Arg Glu Thr Ser Phe Ile His Ala Ile Ser Ser Ala Gly Val
                85   90   95
Met Tyr Ile Ile Thr Lys Asn Cys Ser Met Gly Asp Phe Glu Asn Cys
            100   105   110
Gly Cys Asp Gly Ser Asn Asn Gly Lys Thr Gly Gly His Gly Trp Ile
        115   120   125
Trp Gly Gly Cys Ser Asp Asn Val Glu Phe Gly Glu Arg Ile Ser Lys
    130   135   140
Leu Phe Val Asp Ser Leu Glu Lys Gly Lys Asp Ala Arg Ala Leu Met
145   150   155   160
Asn Leu His Asn Asn Arg Ala Gly Arg Leu Ala Val Arg Ala Thr Met
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Lys Arg Thr Cys Lys Cys His Gly Ile Ser Gly Ser Cys Ser Ile Gln
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Thr Cys Trp Leu Gln Leu Ala Glu Phe Arg Glu Met Gly Asp Tyr Leu
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Phe Leu Pro Ser Ala Glu Ala Glu Leu Ile Phe Leu Glu Glu Ser Pro
                245                 250                 255
Asp Tyr Cys Thr Cys Asn Ser Ser Leu Gly Ile Tyr Gly Thr Glu Gly
            260                 265                 270
Arg Glu Cys Leu Gln Asn Ser His Asn Thr Ser Arg Trp Glu Arg Arg
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Ser Cys Gly Arg Leu Cys Thr Glu Cys Gly Leu Gln Val Glu Glu Arg
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Lys Thr Glu Val Ile Ser Ser Cys Asn Cys Lys Phe Gln Trp Cys Cys
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<213> homo sapiens

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<210> 21
<211> 34
<212> PRT
<213> homo sapiens

<400> 21
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Pro Ile
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<210> 22
<211> 111
<212> DNA
<213> homo sapiens

<400> 22
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<210> 23
<211> 36
<212> PRT
<213> homo sapiens

<400> 23

Met Gly His Leu Val Leu His Gly His Pro Gly Thr Leu Gln Val Glu
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            20   25   30
Pro Lys Val Gly
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<210> 24
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<212> DNA
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